Author Index

Abe, K., Yamashita, H., Arai, S. and Kurihara, Y.
Molecular cloning of curculin, a novel taste-modifying protein with a sweet taste (1130) 232
Åkerström, B., see Lindqvist, A. (1130) 63
Albrandt, K., Mull, E., Cooper, G.J.S. and Johnson, M.J.
Nucleotide sequence of a cDNA for canine amylin (1130) 97
Altieri, M., see Lindqvist, A. (1130) 63
Aoyama, Y., see Oka, T. (1130) 182
Appelhans, H., see Gehrig, H. (1130) 326
Arai, S., see Abe, K. (1130) 232
Ariga, H., see Taira, T. (1130) 166
Arndt, E.
The genes for ribosomal protein L15 and the protein equivalent to secY in the archaebacterium *Haloarcula* (*Halobacterium*) *marismortui* (1130) 113
Artiaco, D., see Furia, M. (1130) 314

Bai, G., Nichols, R.A. and Weiss, B.
Cyclic AMP selectively up-regulates calmodulin genes I and II in PC12 cells (1130) 189
Bakhshi, M.R., see Sinha, D. (1130) 209
Bergeron, J.M., see Yang, F. (1130) 149
Bertchtold, M.W., see Furter, C.S. (1130) 213
Biard, D.S.F., James, M.R., Cordier, A. and Sarasin, A.
Regulation of the *Escherichia coli lac* operon expressed in human cells (1130) 68
Bloemendal, H., see Merck, K.B. (1130) 267
Boussios, T., see Lee, K.M. (1130) 343
Boussios, T., see Li, H. (1130) 218
Bradley, A., see Matzuk, M.M. (1130) 105
Bratt, T., see Lindqvist, A. (1130) 63
Breen, G.A.M., see Jordan, E.M. (1130) 122

Campbell, D.G., see Mann, D.J. (1130) 100
Cech, D., see Schmidt, S. (1130) 41
Chen, Z.-L., see Yang, F. (1130) 149
Cohen, P.T.W., see Mann, D.J. (1130) 100
Conley, A.J., Graham-Lorence, S.E., Kagimoto, M., Lorence, M.C., Murry, B.A., Oka, K., Sanders, D. and Mason, J.I.
Nucleotide sequence of a cDNA encoding porcine testis 17α-hydroxylase cytochrome *P*-450 (1130) 75
Cooper, G.J.S., see Albrandt, K. (1130) 97
Cordier, A., see Biard, D.S.F. (1130) 68
Cupples, R.L., see Yang, F. (1130) 149
Czaika, G. and Mamet-Bratley, M.D.
Defective DNA injection by alkylated and nonalkylated bacteriophage T7 (1130) 52

Davie, J.R., see Hendzel, M.J. (1130) 307
De Crombrugghe, B., see Elima, K. (1130) 78
De Haard-Hoekman, W.A., see Merck, K.B. (1130) 267
De Jong, W.W., see Merck, K.B. (1130) 267
Depew, R.E., see Yang, T.-P. (1130) 227
Deuchars, K.L., Duthie, M. and Ling, V.
Identification of distinct P-glycoprotein gene sequences in rat (1130) 157
Digilio, F.A., see Furia, M. (1130) 314
Dimberg, J., Gustafson-Svärd, C., Weström, B., Tagesson, C. and Söderkvist, P.
Group I phospholipase A_2 mRNA expression in rat glandular stomach and pancreas. Ontogenic development and effects of cortisone acetate (1130) 47
Dumbsky, M., see Haas, A. (1130) 81
Duthie, M., see Deuchars, K.L. (1130) 157

Eerola, I., see Elima, K. (1130) 78
Elela, S.A. and Nazar, R.N.
Extended secondary structure as a basis of increased RNA stability in a thermophilic alga *Cyanidium caldarium* (1130) 339
Elima, K., Metsäranta, M., Kallio, J., Perälä, M., Eerola, I., Garofalo, S., De Crombrugghe, B. and Vuorio, E.
Specific hybridization probes for mouse α2(IX) and α1(X) collagen mRNAs (1130) 78
El'skaya, A.V., see Soldatkin, K.O. (1130) 38
Endo, Y., see Oka, T. (1130) 182
Endo, Y., see Oka, T. (1130) 224

Favia, G., see Furia, M. (1130) 314
Feng, J. and Villeponteau, B.
High-resolution analysis of c-*fos* chromatin accessibility using a novel DNase I-PCR assay (1130) 253
Flannigan, D., see Harauz, G. (1130) 289
Francès, V., Morlé, F. and Godet, J.
Identification of two critical base pairings in 5′ untranslated regions affecting translation efficiency of synthetic uncapped globin mRNAs (1130) 29
Friedrichs, W.E., see Yang, F. (1130) 149
Fujimoto, S., see Itoh, S. (1130) 133
Fujinaga, K., see Ohashi, Y. (1130) 175
Furia, M., Digilio, F.A., Artiaco, D., Favia, G. and Polito, L.C.
Molecular characterization of a *Drosophila melanogaster* variant strain defective in the *Sgs-4* gene dosage compensation (1130) 314
Furter, C.S., Rentsch, J.M. and Bertchtold, M.W.
Sequence variations and promoter activities of long terminal repeats from rat intracisternal A-particles (1130) 213

Galley, K.A., Singh, B. and Gupta, R.S.
Cloning of HSP70 (dnaK) gene from *Clostridium perfringens* using a general polymerase chain reaction based approach (1130) 203
Garofalo, S., see Elima, K. (1130) 78
Gehrig, H., Müller, W. and Appelhans, H.
Complete nucleotide sequence of band 3 related anion transport protein AE2 from human kidney (1130) 326
Gick, G., see Liu, B. (1130) 336
Gilbert, H.J., see Hall, J. (1130) 259
Girardet, J.-L. and Ramstein, J.
A reexamination of the stopped-flow measured hydrogen-deuterium exchange rates in nucleic acid double helices (1130) 127
Godet, J., see Francès, V. (1130) 29

Author Index

Abe, K., Yamashita, H., Arai, S. and Kurihara, Y.
Molecular cloning of curculin, a novel taste-modifying protein with a sweet taste (1130) 232
Åkerström, B., see Lindqvist, A. (1130) 63
Albrandt, K., Mull, E., Cooper, G.J.S. and Johnson, M.J.
Nucleotide sequence of a cDNA for canine amylin (1130) 97
Altieri, M., see Lindqvist, A. (1130) 63
Aoyama, Y., see Oka, T. (1130) 182
Appelhans, H., see Gehrig, H. (1130) 326
Arai, S., see Abe, K. (1130) 232
Ariga, H., see Taira, T. (1130) 166
Arndt, E.
The genes for ribosomal protein L15 and the protein equivalent to secY in the archaebacterium *Haloarcula* (*Halobacterium*) *marismortui* (1130) 113
Artiaco, D., see Furia, M. (1130) 314

Bai, G., Nichols, R.A. and Weiss, B.
Cyclic AMP selectively up-regulates calmodulin genes I and II in PC12 cells (1130) 189
Bakhshi, M.R., see Sinha, D. (1130) 209
Bergeron, J.M., see Yang, F. (1130) 149
Bertchtold, M.W., see Furter, C.S. (1130) 213
Biard, D.S.F., James, M.R., Cordier, A. and Sarasin, A.
Regulation of the *Escherichia coli lac* operon expressed in human cells (1130) 68
Bloemendal, H., see Merck, K.B. (1130) 267
Boussios, T., see Lee, K.M. (1130) 343
Boussios, T., see Li, H. (1130) 218
Bradley, A., see Matzuk, M.M. (1130) 105
Bratt, T., see Lindqvist, A. (1130) 63
Breen, G.A.M., see Jordan, E.M. (1130) 122

Campbell, D.G., see Mann, D.J. (1130) 100
Cech, D., see Schmidt, S. (1130) 41
Chen, Z.-L., see Yang, F. (1130) 149
Cohen, P.T.W., see Mann, D.J. (1130) 100
Conley, A.J., Graham-Lorence, S.E., Kagimoto, M., Lorence, M.C., Murry, B.A., Oka, K., Sanders, D. and Mason, J.I.
Nucleotide sequence of a cDNA encoding porcine testis 17α-hydroxylase cytochrome *P*-450 (1130) 75
Cooper, G.J.S., see Albrandt, K. (1130) 97
Cordier, A., see Biard, D.S.F. (1130) 68
Cupples, R.L., see Yang, F. (1130) 149
Czaika, G. and Mamet-Bratley, M.D.
Defective DNA injection by alkylated and nonalkylated bacteriophage T7 (1130) 52

Davie, J.R., see Hendzel, M.J. (1130) 307
De Crombrugghe, B., see Elima, K. (1130) 78
De Haard-Hoekman, W.A., see Merck, K.B. (1130) 267
De Jong, W.W., see Merck, K.B. (1130) 267
Depew, R.E., see Yang, T.-P. (1130) 227
Deuchars, K.L., Duthie, M. and Ling, V.
Identification of distinct P-glycoprotein gene sequences in rat (1130) 157
Digilio, F.A., see Furia, M. (1130) 314
Dimberg, J., Gustafson-Svärd, C., Weström, B., Tagesson, C. and Söderkvist, P.
Group I phospholipase A_2 mRNA expression in rat glandular stomach and pancreas. Ontogenic development and effects of cortisone acetate (1130) 47
Dumbsky, M., see Haas, A. (1130) 81
Duthie, M., see Deuchars, K.L. (1130) 157

Eerola, I., see Elima, K. (1130) 78
Elela, S.A. and Nazar, R.N.
Extended secondary structure as a basis of increased RNA stability in a thermophilic alga *Cyanidium caldarium* (1130) 339
Elima, K., Metsäranta, M., Kallio, J., Perälä, M., Eerola, I., Garofalo, S., De Crombrugghe, B. and Vuorio, E.
Specific hybridization probes for mouse α2(IX) and α1(X) collagen mRNAs (1130) 78
El'skaya, A.V., see Soldatkin, K.O. (1130) 38
Endo, Y., see Oka, T. (1130) 182
Endo, Y., see Oka, T. (1130) 224

Favia, G., see Furia, M. (1130) 314
Feng, J. and Villeponteau, B.
High-resolution analysis of c-*fos* chromatin accessibility using a novel DNase I-PCR assay (1130) 253
Flannigan, D., see Harauz, G. (1130) 289
Francès, V., Morlé, F. and Godet, J.
Identification of two critical base pairings in 5′ untranslated regions affecting translation efficiency of synthetic uncapped globin mRNAs (1130) 29
Friedrichs, W.E., see Yang, F. (1130) 149
Fujimoto, S., see Itoh, S. (1130) 133
Fujinaga, K., see Ohashi, Y. (1130) 175
Furia, M., Digilio, F.A., Artiaco, D., Favia, G. and Polito, L.C.
Molecular characterization of a *Drosophila melanogaster* variant strain defective in the *Sgs-4* gene dosage compensation (1130) 314
Furter, C.S., Rentsch, J.M. and Bertchtold, M.W.
Sequence variations and promoter activities of long terminal repeats from rat intracisternal A-particles (1130) 213

Galley, K.A., Singh, B. and Gupta, R.S.
Cloning of HSP70 (dnaK) gene from *Clostridium perfringens* using a general polymerase chain reaction based approach (1130) 203
Garofalo, S., see Elima, K. (1130) 78
Gehrig, H., Müller, W. and Appelhans, H.
Complete nucleotide sequence of band 3 related anion transport protein AE2 from human kidney (1130) 326
Gick, G., see Liu, B. (1130) 336
Gilbert, H.J., see Hall, J. (1130) 259
Girardet, J.-L. and Ramstein, J.
A reexamination of the stopped-flow measured hydrogen-deuterium exchange rates in nucleic acid double helices (1130) 127
Godet, J., see Francès, V. (1130) 29

Gor, D. and Mayfield, J.E.
Cloning and nucleotide sequence of the *Brucella abortus gro*E operon (1130) 120
Graham-Lorence, S.E., see Conley, A.J. (1130) 75
Gupta, R.S., see Galley, K.A. (1130) 203
Gupta, R.S., see Rusanganwa, E. (1130) 90
Gustafson-Svärd, C., see Dimberg, J. (1130) 47

Haas, A., Dumbsky, M. and Kreft, J.
Listeriolysin genes: complete sequence of *ilo* from *Listeria ivanovii* and of *lso* from *Listeria seeligeri* (1130) 81
Hall, J., Hirst, B.H., Hazlewood, G.P. and Gilbert, H.J.
The use of chimeric gene constructs to express a bacterial endoglucanase in mammalian cells (1130) 259
Handa, H. and Nakajima, K.
The gene for $tRNA^{Lys}$ is encoded in the rapeseed (*Brassica napus L.*) mitochondrial DNA (1130) 117
Harauz, G. and Flannigan, D.
Characteristic electron microscopical projections of the small ribosomal subunit from *Thermomyces lanuginosus* (1130) 289
Hashimoto, H., see Itoh, S. (1130) 133
Hashimoto, T., see Tomioka, Y. (1130) 109
Hazlewood, G.P., see Hall, J. (1130) 259
Heintz, N.H., see Held, P.G. (1130) 235
Held, P.G. and Heintz, N.H.
Eukaryotic replication origins (1130) 235
Helfrich, W., see Rossell, R.J. (1130) 95
Hendzel, M.J. and Davie, J.R.
Nuclear distribution of histone deacetylase: a marker enzyme for the internal nuclear matrix (1130) 307
Hirose, A., see Tomioka, Y. (1130) 109
Hirst, B.H., see Hall, J. (1130) 259
Hishinuma, T., see Tomioka, Y. (1130) 109
Hsie, D.Y., see Rabek, J.P. (1130) 317
Hudspeth, D.S.S. and Vary, P.S.
spoVG sequence of *Bacillus megaterium* and *Bacillus subtilis* (1130) 229

Ichimura, S., see Nenoi, M. (1130) 247
Iguchi-Ariga, S.M.M., see Taira, T. (1130) 166
Iizasa, T., see Ishizuka, T. (1130) 139
Inoue, A., Yanagisawa, M. and Masaki, T.
Differential tissue expression of multiple genes for chicken smooth muscle/nonmuscle myosin regulatory light chains (1130) 197
Ishijima, S., see Ishizuka, T. (1130) 139
Ishikura, H., see Matsugi, J. (1130) 333
Ishizuka, T., Iizasa, T., Taira, M., Ishijima, S., Sonoda, T., Shimada, H., Nagatake, N. and Tatibana, M.
Promoter regions of the human X-linked housekeeping genes *PRPS1* and *PRPS2* encoding phosphoribosylpyrophosphate synthetase subunit I and II isoforms (1130) 139
Ito, M., see Oka, T. (1130) 224
Itoh, S., Yanagimoto, T., Tagawa, S., Hashimoto, H., Kitamura, R., Nakajima, Y., Okochi, T., Fujimoto, S., Uchino, J. and Kamataki, T.
Genomic organization of human fetal specific *P*-450IIIA7 (cytochrome *P*-450HFLa)-related gene(s) and interaction of transcriptional regulatory factor with its DNA element in the 5′ flanking region (1130) 133
Itoh, S., see Yanagimoto, T. (1130) 329

James, M.R., see Biard, D.S.F. (1130) 68
Jia, H.-T., see Matsugi, J. (1130) 333
Johnson, M.J., see Albrandt, K. (1130) 97
Jordan, E.M. and Breen, G.A.M.
Molecular cloning of an import precursor of the δ-subunit of the human mitochondrial ATP synthase complex (1130) 122

Kagimoto, M., see Conley, A.J. (1130) 75
Kallio, J., see Elima, K. (1130) 78
Kamataki, T., see Itoh, S. (1130) 133
Kamataki, T., see Yanagimoto, T. (1130) 329
Kastern, W., see Lindqvist, A. (1130) 63
Katano, T., see Oka, T. (1130) 182
Kemshead, J.T., see Rossell, R.J. (1130) 95
Kihara, F., see Taira, T. (1130) 166
Kirby, E.P., see Sinha, D. (1130) 209
Kitamura, R., see Itoh, S. (1130) 133
Kodaira, K.-I., Nakano, K., Okada, S. and Taketo, A.
Nucleotide sequence of the genome of the bacteriophage $\alpha 3$: interrelationship of the genome structure and the gene products with those of the phages, ϕX174, G4 and ϕK (1130) 277
Koval'chuke, O., see Soldatkin, K.O. (1130) 38
Kreft, J., see Haas, A. (1130) 81
Krynetskaya, N.F., see Schmidt, S. (1130) 41
Kurihara, Y., see Abe, K. (1130) 232

Lazarus, P., see Ronai, Z. (1130) 20
Lee, K.M., Subar, M., Li, H. and Boussios, T.
Cloning of two adult hamster globin cDNAs (α and β^{major}) (1130) 343
Lee, K.M., see Li, H. (1130) 218
Li, H., Subar, M., Lee, K.M. and Boussios, T.
Cloning and sequence analysis of two embryonic β-like globin cDNAs (y and z) of hamster (1130) 218
Li, H., see Lee, K.M. (1130) 343
Lindqvist, A., Bratt, T., Altieri, M., Kastern, W. and Åkerström, B.
Rat α_1-microglobulin: co-expression in liver with the light chain of inter-α-trypsin inhibitor (1130) 63
Ling, V., see Deuchars, K.L. (1130) 157
Liu, B. and Gick, G.
Characterization of the 5′ flanking region of the rat Na^+/K^+-ATPase $\beta 1$ subunit gene (1130) 336
Lorence, M.C., see Conley, A.J. (1130) 75
Lorens, J.B., see Male, R. (1130) 345
Lossius, I., see Male, R. (1130) 345

Male, R., Nerland, A.H., Lorens, J.B., Telle, W., Lossius, I. and Totland, G.K.
The complete nucleotide sequence of the Atlantic salmon growth hormone I gene (1130) 345
Malik, A., see Rossell, R.J. (1130) 95
Mamet-Bratley, M.D., see Czaika, G. (1130) 52
Mann, D.J., Campbell, D.G., McGowan, C.H. and Cohen, P.T.W.
Mammalian protein serine/threonine phosphatase 2C: cDNA cloning and comparative analysis of amino acid sequences (1130) 100
Masaki, T., see Inoue, A. (1130) 197
Mason, J.I., see Conley, A.J. (1130) 75
Matsugi, J., Jia, H.-T., Murao, K. and Ishikura, H.
Nucleotide sequences of serine tRNAs from *Bacillus subtilis* (1130) 333
Matzuk, M.M. and Bradley, A.
Cloning of the human activin receptor cDNA reveals high evolutionary conservation (1130) 105
Mayer, D., see Schiebel, K. (1130) 349
Mayfield, J.E., see Gor, D. (1130) 120
McCarthy, J.E.G., see Ziemke, P. (1130) 297
McGowan, C.H., see Mann, D.J. (1130) 100
Merck, K.B., De Haard-Hoekman, W.A., Oude Essink, B.B., Bloemendal, H. and De Jong, W.W.

Expression and aggregation of recombinant αA-crystallin and its two domains (1130) 267
Metelev, V.G., see Schmidt, S. (1130) 41
Metsäranta, M., see Elima, K. (1130) 78
Mita, K., see Nenoi, M. (1130) 247
Miyamoto, K.-i., see Oka, T. (1130) 224
Mizugaki, M., see Tomioka, Y. (1130) 109
Moritani, H., see Tomioka, Y. (1130) 109
Moriuchi, T., see Ohashi, Y. (1130) 175
Morlé, F., see Francès, V. (1130) 29
Mull, E., see Albrandt, K. (1130) 97
Muller-Enoch, D., see Yanagimoto, T. (1130) 329
Müller, W., see Gehrig, H. (1130) 326
Murao, K., see Matsugi, J. (1130) 333
Murry, B.A., see Conley, A.J. (1130) 75

Nagatake, N., see Ishizuka, T. (1130) 139
Nakajima, K., see Handa, H. (1130) 117
Nakajima, Y., see Itoh, S. (1130) 133
Nakano, K., see Kodaira, K.-I. (1130) 277
Natori, Y., see Oka, T. (1130) 182
Natori, Y., see Oka, T. (1130) 224
Nazar, R.N., see Elela, S.A. (1130) 339
Negishi, Y., see Taira, T. (1130) 166
Nenoi, M., Mita, K. and Ichimura, S.
Evolutionarily conserved structure of the 3′ non-translated region of a Chinese hamster polyubiquitin gene (1130) 247
Nerland, A.H., see Male, R. (1130) 345
Nichols, R.A., see Bai, G. (1130) 189
Niemann, A., see Schmidt, S. (1130) 41

Ohashi, Y., Sawada, Y., Moriuchi, T. and Fujinaga, K.
Analysis of the 5′ flanking region of the rat proliferating cell nuclear antigen (PCNA) gene (1130) 175
Oka, K., see Conley, A.J. (1130) 75
Oka, T., Aoyama, Y., Natori, Y., Katano, T. and Endo, Y.
An efficient expression system for a variant form of the cytotoxic protein α-sarcin in *Escherichia coli* (1130) 182
Oka, T., Endo, Y., Ito, M., Miyamoto, K.-i., Sasakawa, T., Suzuki, I. and Natori, Y.
Molecular cloning of chick liver HMG 2a cDNA and developmental expression of HMG 2a mRNA (1130) 224
Okada, S., see Kodaira, K.-I. (1130) 277
Okochi, T., see Itoh, S. (1130) 133
Oretskaya, T.S., see Schmidt, S. (1130) 41
Oude Essink, B.B., see Merck, K.B. (1130) 267

Papaconstantinou, J., see Rabek, J.P. (1130) 317
Patel, K., see Rossell, R.J. (1130) 95
Pegg, A.E., see Tekwani, B.L. (1130) 221
Pekel, E., see Schiebel, K. (1130) 349
Perälä, M., see Elima, K. (1130) 78
Polito, L.C., see Furia, M. (1130) 314
Potapov, A.P., see Soldatkin, K.O. (1130) 38

Rabek, J.P., Hsie, D.Y. and Papaconstantinou, J.
α-Fetoprotein expression in fetal kidney cells does not require enhancers (1130) 317
Ramstein, J., see Girardet, J.-L. (1130) 127
Rentsch, J.M., see Furter, C.S. (1130) 213
Robbins, S.M., Williams, J.G., Spiegelman, G.B. and Weeks, G.
Cloning and characterization of the *Dictyostelium discoideum rasG* genomic sequences (1130) 85
Robinson, R., see Ronai, Z. (1130) 20
Ronai, Z., Robinson, R., Rutberg, S., Lazarus, P. and Sardana, M.
Aldolase–DNA interactions in a SEWA cell system (1130) 20
Rossell, R.J., Malik, A., Helfrich, W., Kemshead, J.T. and Patel, K.
A discordance between the human muscle NCAM sequence and those seen in other NCAM cDNA clones (1130) 95
Rusanganwa, E., Singh, B. and Gupta, R.S.
Cloning of HSP60 (*GroEL*) operon from *Clostridium perfringens* using a polymerase chain reaction based approach (1130) 90
Rutberg, S., see Ronai, Z. (1130) 20

Sanders, D., see Conley, A.J. (1130) 75
Sarasin, A., see Biard, D.S.F. (1130) 68
Sardana, M., see Ronai, Z. (1130) 20
Sasakawa, T., see Oka, T. (1130) 224
Sawada, Y., see Ohashi, Y. (1130) 175
Schiebel, K., Pekel, E. and Mayer, D.
The nucleotide sequence of rat liver glycogen phosphorylase cDNA (1130) 349
Schmidt, S., Niemann, A., Krynetskaya, N.F., Oretskaya, T.S., Metelev, V.G., Suchomlinov, V.V., Shabarova, Z.A. and Cech, D.
The use of oligonucleotide probes containing 2′-deoxy-2′-fluoronucleosides for regiospecific cleavage of RNA by RNase H from *Escherichia coli* (1130) 41
Shabarova, Z.A., see Schmidt, S. (1130) 41
Shimada, H., see Ishizuka, T. (1130) 139
Singh, B., see Galley, K.A. (1130) 203
Singh, B., see Rusanganwa, E. (1130) 90
Sinha, D., Bakhshi, M.R. and Kirby, E.P.
Complete cDNA sequence of bovine α1-antitrypsin (1130) 209
Söderkvist, P., see Dimberg, J. (1130) 47
Soldatkin, K.O., Koval'chuke, O., Potapov, A.P. and El'skaya, A.V.
A codon sugar structure strongly influences tRNA binding to programmed ribosomes (1130) 38
Sonoda, T., see Ishizuka, T. (1130) 139
Spiegelman, G.B., see Robbins, S.M. (1130) 85
Stanley, B.A., see Tekwani, B.L. (1130) 221
Subar, M., see Lee, K.M. (1130) 343
Subar, M., see Li, H. (1130) 218
Suchomlinov, V.V., see Schmidt, S. (1130) 41
Suzuki, I., see Oka, T. (1130) 224

Tagawa, S., see Itoh, S. (1130) 133
Tagesson, C., see Dimberg, J. (1130) 47
Taira, M., see Ishizuka, T. (1130) 139
Taira, T., Negishi, Y., Kihara, F., Iguchi-Ariga, S.M.M. and Ariga, H.
c-*myc* protein complex binds to two sites in human hsp70 promoter region (1130) 166
Taketo, A., see Kodaira, K.-I. (1130) 277
Tatibana, M., see Ishizuka, T. (1130) 139
Tekwani, B.L., Stanley, B.A. and Pegg, A.E.
Nucleotide sequence of hamster *S*-adenosylmethionine decarboxylase cDNA (1130) 221
Telle, W., see Male, R. (1130) 345
Thoma, F.
Nucleosome positioning (1130) 1
Tomioka, Y., Hirose, A., Moritani, H., Hishinuma, T., Hashimoto, T. and Mizugaki, M.
cDNA cloning of mitochondrial Δ^3, Δ^2-enoyl-CoA isomerase of rat liver (1130) 109
Totland, G.K., see Male, R. (1130) 345

Uchino, J., see Itoh, S. (1130) 133

Vary, P.S., see Hudspeth, D.S.S. (1130) 229
Villeponteau, B., see Feng, J. (1130) 253
Vuorio, E., see Elima, K. (1130) 78

Weeks, G., see Robbins, S.M. (1130) 85
Weiss, B., see Bai, G. (1130) 189
Weström, B., see Dimberg, J. (1130) 47
Williams, J.G., see Robbins, S.M. (1130) 85

Yamashita, H., see Abe, K. (1130) 232
Yanagimoto, T., Itoh, S., Muller-Enoch, D. and Kamataki, T.
Mouse liver cytochrome *P*-450 (P-450IIIAM1): its cDNA cloning and inducibility by dexamethasone (1130) 329
Yanagimoto, T., see Itoh, S. (1130) 133
Yanagisawa, M., see Inoue, A. (1130) 197
Yang, F., Chen, Z.-L., Bergeron, J.M., Cupples, R.L. and Friedrichs, W.E.
Human α_2-HS-glycoprotein/bovine fetuin homologue in mice: identification and developmental regulation of the gene (1130) 149
Yang, T.-P. and Depew, R.E.
Nucleotide sequence of a region duplicated in *Escherichia coli toc* mutants (1130) 227

Ziemke, P. and McCarthy, J.E.G.
The control of mRNA stability in *Escherichia coli*: manipulation of the degradation pathway of the polycistronic *atp* mRNA (1130) 297